GHJTVIVT HJTVIVTJ

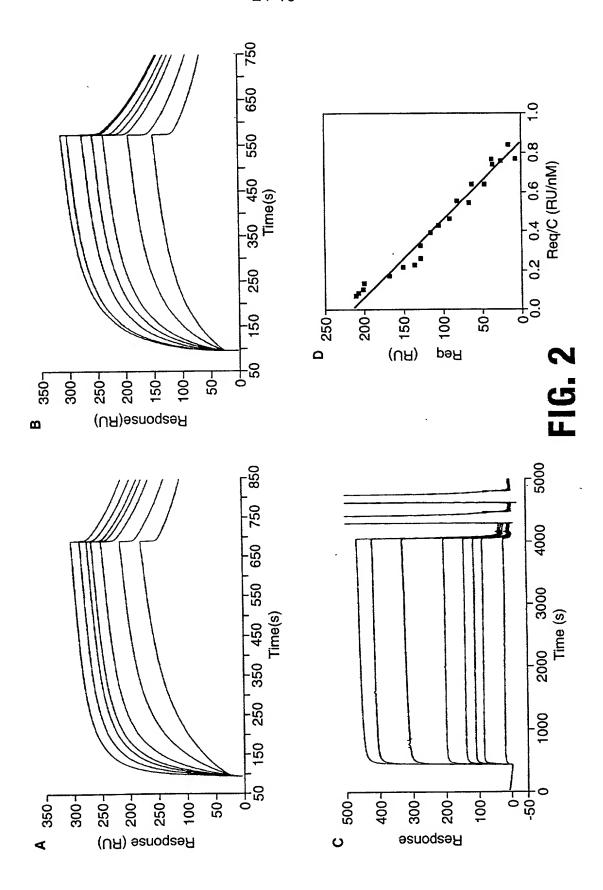
GLTVIVTL DCLTVIVT KDCLTVIV

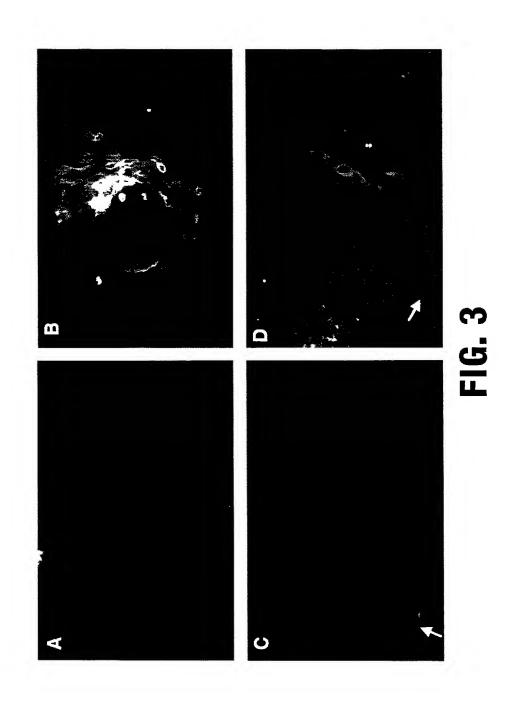
TAAA9DAM

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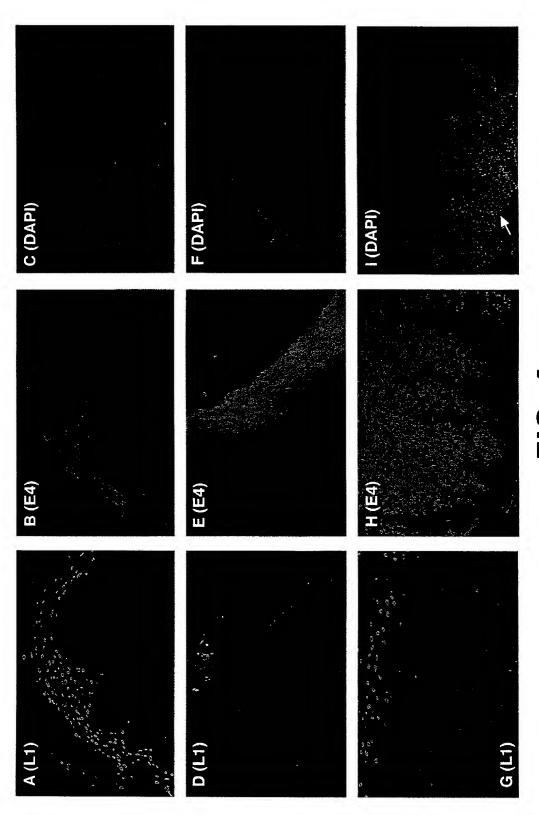


FIG. 4

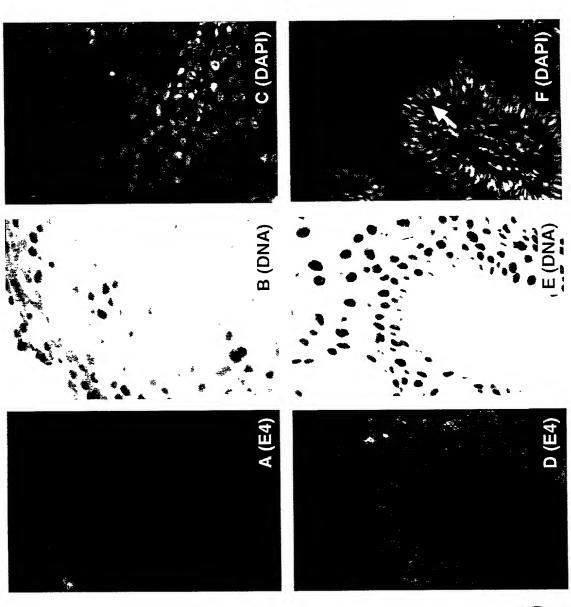
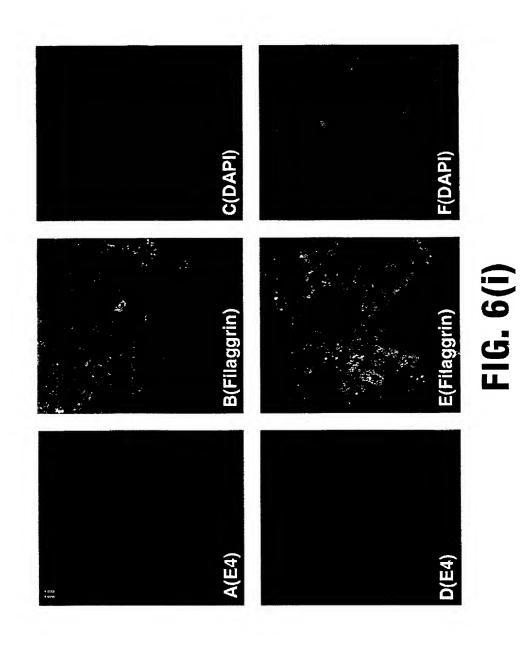


FIG. 5



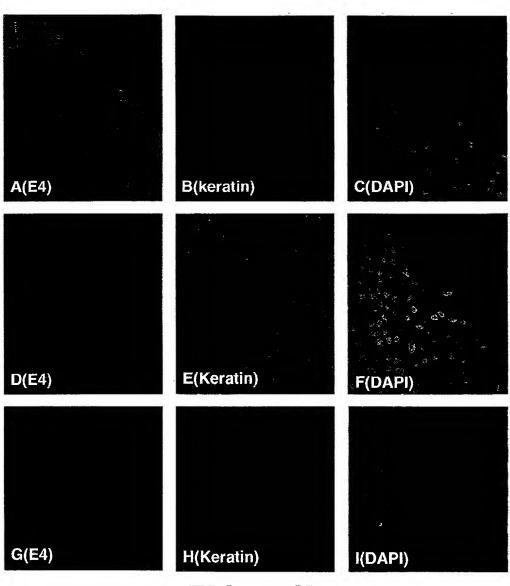


FIG. 6(ii)

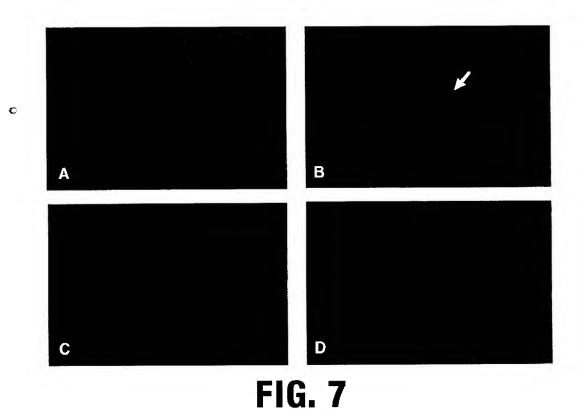


FIG. 8

	CHARGED REGION	
most-likely	PPPPP.RPWAGPKKPTRGPPRRRRLESDS	- 1
HPV54		51 43
HPV32		61
HPV42 HPV3	LT.TTTQTVOTEQHT-CSKPH-HN-T	59
HPV28		46 46
HPV10	K-RWARDRN-SDSDSSTDT-	46
HPV29		52
HPV61 HPV2a		49
HPV27	QEEQLRKRC-P-RRQRVPSAV EQEQLR-Q.TCC-P-RRHRVPSAG	61 58
HPV57		53
HPV26		45
HPV51 HPV30	LPAHNSN-X	42
HPV53	RPPHGRNVLRPPHGRNVL	46 44
HPV56	T-T- OY DOEN	42
HPV66	RYPTDOEN	42
HPV18	ACQRAALH-L	44
HPV45 HPV39	KHQNSLLL	44 43
HPV70		43
HPV59		44
HPV7		45
HPV40 HPV16	S.A-P-H-PE	48 41
HPV35h		43
HPV31	······	46
HPV52		40
HPV33 HPV58	HH-QPNDD-LTTKVH-GOD.	30
RhPV1	PR-TQECQEC	29 43
HVP6b		40
HPV11	·····. AC GEH	41
HVP44 HVP55		48
HVP13	······································	48 43
PCPV1	······················	44
HPV34 HPV19	THRTRVCOHGNGT	35
HPV25	GTRDDLPAGPDDKKARN.DQG-NPSP-RG-G-GFRLTGDHDPNPPAGHDDSKPKRARGDQG-S-GGGP.SPA-VSDRG-G-GNLRLSGDQDP	93
HPV20	GTDGDLPVGQGEQ-KARGDGPGQSPSPS-G-G-GRGTGLGLGLGL	105 91
HPV21	·····SP-RG-G-GR-LG-PDPGPDP	90
HPV14d HPV5	EGTDADRPVGPGER-KGRG-DRGSP-RG-G-GGLDPGRNR	92
HPV36	SQGDRKRSKGDQGPDTGPGLGPGRG-S-KTLGP-PG-GPR-S-RLGPLQADRDPEEGQGDRKRSKGDQGPDTDPLGPDRG-S-GT-Q.PL-LPPGLR-SPGSGYQPD	115 114
HPV47	OGDRKRTKGDPDPDPGRGPVLKPTLPTGLRRSTRLVLV-GOGPPP D LPA	116
HPV12	GDRKRSKGDOGRDTAPSLTPGRA-S-KGL-P-PYGPPR-SHG. TGGRD	107
HPV8 HPV24	QDRKKSRGDQGRDTAPGLAPGRS-GL.GL-P-PYGPR-SP-QFGPGPDRDPE	111
HPV15	PLTPDADDDPRPGKRSKGDEHG-A-GAA-PL-LDL.DPPQ-G-DQPPGATGG TTEKNLALQPGGKD.KDKDTQ.QGDQQGGDKKPG	102 86
HPV17	DTGGKRLALQPGTKD-TSDDQHGGDKQPG	79
HPV37	EEKHLALQGKKDKE-T-QQGDQPGGNKQPPG	82
HPV9	PPPGRKDRDKEKEKEKEKEK.KTT-D-GDP.RVEOKPKGEG-	93
HPV22 HPV23	LVLQSP-SGGKKGE.RDKDQQGEEK-DQGPAP	74
HPV38	KHLALQG.GK.KDKESPGEEK-DQGPG.AE DTGEKHLALQAG.KGKD-EKPQA-KGEEKADQG	71 71
HPV49	.TLVLQQPPTPGKRSRDDDPGLEPG-ADGKAPQAVPD-DPDPLPDPEGP	71 120
HPV4	P-P-LKPREYDY-E	57
HPV65 HPV48	SL-RALVVG-NRGNLNRPPQR-PKP-GYY-E	57
HPV50	LE.GDASQKTPTSP-HPDYE ANRKD.LEAV.NQYR-PNH-P-HQQY-F	49 50
HPV60	LTEDR-H-RESLALVFDYDAE	55
BPV1		45
BPV2 EEPV	YPSOA	44
DPV		44 46
BPV4	S-LRD-DHGHDH.DRL	50
HPV41		38
COPV	LGK.GRHG-LDGGRGSPEGODEE	48
CRPV ROPV	QG-KVHDEGQGHQGC.NEG LOY-Q.ART.IRRSSRY-G-FVT-GGDPDPQ	51 55
HVP1a		55 50
HPV63	KL-EKQ-RRGRDTTRNLFAG	46
MnPV	·····GGNQQRGRGA	59

FIG. 9

.... CHARGED REGION -

most-likely DSDSGEVEG PTPTTPPAPPTG HPV54 E-TA.QTSN H-APQT HPV32 --LC.SHQQ S-CS-T.-SQ-Y HPV42 --VDSRHHS TCS-QT--S-ASP HPV3 N-S-NSNSN NI-KP--RK-LN 72 57 80 81 67 60 HPV28 61 HPV10 73 HPV29 68 HPV61 79 HPV2a S--- SI SG--LRERSER- S--- S- NS--LRGRSEK- S--- S- NS--LRGRSEK- S--- S--- NS--LRGRSEK- S--- NS--- NS--LRGRSEK- S--- NS--- NS--LRGRSEK- S--- NS--- NS----76 HPV27 HPV57 71 HPV26 65 HPV30 66 HPV53 63 HPV56 62 HPV66 62 53 HPV18 55 HPV45 HPV39 56 51 HPV70 HPV59 67 HPV7 HPV40 59 HPV16 58 HPV35h 64 HPV31 HPV52 58 HPV33 45 HPV58 49 RhPV1 55 HVP6b HPV11 55 HVP44 HVP55 HVP13 PCPV1 62 HPV34 41 HPV19 119 HPV25 149 HPV20 144 HPV21 152 HPV14d 135 HPV5 147 HPV36 144 HPV47 145 HPV12 129 HPV8 147 HPV24 120 HPV15 HPV17 103 HPV37 106 HPV9 112 HPV22 95 HPV23 92 99 HPV38 140 HPV49 HPV4 77 HPV65 77 HPV48 67 HPV 50 70 HPV60 75 V.GY-T-LA R---IF.LQARV.GY-T-LA R---IF.LQARE.LAKTGV-P F-ARL-T-HHHP BPV1 BPV2 EEPV DPV BPV4 71 HPV41 COPV 63 CRPV ROPV HVP1aPT-E...-.....G-EV-EI--SD..... 62 NPV63 MnPVRTP-P-TTA.....QR-KR-RRAC-RK.... 81

FIG. 9 CONT'D